

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/535,522
Source: Pg 1/10
Date Processed by STIC: 5/27/05

ENTERED

Raw Sequence Listing before editing,
for reference only



PCT

RAW SEQUENCE LISTING

DATE: 05/27/2005

PATENT APPLICATION: US/10/535,522

TIME: 11:22:32

Input Set : A:\PTO.DA.txt

Output Set: N:\CRF4\05272005\J535522.raw

3 <110> APPLICANT: Steinkasserer, Alexander
 5 <120> TITLE OF INVENTION: Use of Soluble Forms of CD83 and Nucleic Acids Encoding
 6 them for the Treatment or Prevention of Diseases
 W--> 7 <130> FILE REFERENCE: 032723woJH
 W--> 8 <140> CURRENT APPLICATION NUMBER:
 C--> 9 <141> CURRENT FILING DATE: 2005-05-18
 W--> 10 <160> NUMBER OF SEQ ID: 12
 11 <170> SOFTWARE: PatentIn Ver. 2.1

ERRORED SEQUENCES

**Does Not Comply
 Corrected Diskette Needed**

356 <210> SEQ ID NO: 8
 357 <211> LENGTH: 139
 358 <212> TYPE: PRT
 359 <213> ORGANISM: Artificial Sequence
 W--> 360 <220> FEATURE: *insert <220>*
 360 <223> OTHER INFORMATION: Description of Artificial Sequence: partial
 361 sequence of pGEX2ThCD83ext
 OK> 363 <400> SEQUENCE: 8
 364 Pro Pro Lys Ser Asp Leu Val Pro Arg Gly Ser Pro Gly Thr Pro Glu
 365 -5 -1 1 5
 367 Val Lys Val Ala Cys Ser Glu Asp Val Asp Leu Pro Cys Thr Ala Pro
 368 10 15 20
 370 Trp Asp Pro Gln Val Pro Tyr Thr Val Ser Trp Val Lys Leu Leu Glu
 371 25 30 35
 373 Gly Gly Glu Glu Arg Met Glu Thr Pro Gln Glu Asp His Leu Arg Gly
 374 40 45 50 55
 376 Gln His Tyr His Gln Lys Gly Gln Asn Gly Ser Phe Asp Ala Pro Asn
 377 60 65 70
 379 Glu Arg Pro Tyr Ser Leu Lys Ile Arg Asn Thr Thr Ser Cys Asn Ser
 380 75 80 85
 382 Gly Thr Tyr Arg Cys Thr Leu Gln Asp Pro Asp Gly Gln Arg Asn Leu
 383 90 95 100
 385 Ser Gly Lys Val Ile Leu Arg Val Thr Gly Cys Pro Ala Gln Arg Lys
 386 105 110 115
 388 Glu Glu Thr Phe Lys Lys Tyr Arg Ala Glu Ile
 389 120 125 130
 444 <210> SEQ ID NO: 10
 445 <211> LENGTH: 139
 446 <212> TYPE: PRT
 447 <213> ORGANISM: Artificial Sequence
 W--> 448 <220> FEATURE: *insert <220>*

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/535,522

DATE: 05/27/2005

TIME: 11:22:32

Input Set : A:\PTO.DA.txt

Output Set : N:\CRF4\05272005\J535522.raw

448 <223> OTHER INFORMATION: Description of Artificial Sequence: partial
 449 sequence of pGEX2ThCD83ext_mut129_CtoS

451 <400> SEQUENCE: 11/10

452 Pro Pro Lys Ser Asp Leu Val Pro Arg Gly Ser Pro Gly Thr Pro Glu
 453 -5 -1 1 5
 455 Val Lys Val Ala Cys Ser Glu Asp Val Asp Leu Pro Cys Thr Ala Pro
 456 10 15 20
 458 Trp Asp Pro Gln Val Pro Tyr Thr Val Ser Trp Val Lys Leu Leu Glu
 459 25 30 35
 461 Gly Gly Glu Glu Arg Met Glu Thr Pro Gln Glu Asp His Leu Arg Gly
 462 40 45 50 55
 464 Gln His Tyr His Gln Lys Gly Gln Asn Gly Ser Phe Asp Ala Pro Asn
 465 60 65 70
 467 Glu Arg Pro Tyr Ser Leu Lys Ile Arg Asn Thr Thr Ser Cys Asn Ser
 468 75 80 85
 470 Gly Thr Tyr Arg Cys Thr Leu Gln Asp Pro Asp Gly Gln Arg Asn Leu
 471 90 95 100
 473 Ser Gly Lys Val Ile Leu Arg Val Thr Gly Ser Pro Ala Gln Arg Lys
 474 105 110 115
 476 Glu Glu Thr Phe Lys Lys Tyr Arg Ala Glu Ile
 478 120 125 130

494 <210> SEQ ID NO: 12

495 <211> LENGTH: 66

496 <212> TYPE: DNA

497 <213> ORGANISM: Artificial Sequence

W--> 498 <220> FEATURE:

499 <223> OTHER INFORMATION: Description of Artificial Sequence: primer
 500 antisense-CD83extra_mutantCtoS

502 <400> SEQUENCE: 12

503 aattagaatt ctcaaatctc cgctctgtat ttcttaaaag tctcttcttt acgctgtgca 60
 504 ggggat 66

E--> 509 2 -

VERIFICATION SUMMARY

DATE: 05/27/2005

PATENT APPLICATION: US/10/535,522

TIME: 11:22:33

Input Set : A:\PTO.DA.txt

Output Set: N:\CRF4\05272005\J535522.raw

L:7 M:283 W: Missing Blank Line separator, <130> field identifier
L:8 M:283 W: Missing Blank Line separator, <140> field identifier
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:10 M:283 W: Missing Blank Line separator, <160> field identifier
L:18 M:283 W: Missing Blank Line separator, <220> field identifier
L:128 M:283 W: Missing Blank Line separator, <220> field identifier
L:286 M:283 W: Missing Blank Line separator, <220> field identifier
L:297 M:283 W: Missing Blank Line separator, <220> field identifier
L:308 M:283 W: Missing Blank Line separator, <220> field identifier
L:311 M:283 W: Missing Blank Line separator, <220> field identifier
L:314 M:283 W: Missing Blank Line separator, <220> field identifier
L:360 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8
L:363 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:8
L:396 M:283 W: Missing Blank Line separator, <220> field identifier
L:399 M:283 W: Missing Blank Line separator, <220> field identifier
L:402 M:283 W: Missing Blank Line separator, <220> field identifier
L:448 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:10
L:451 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:10
L:451 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:10 differs:11
L:486 M:283 W: Missing Blank Line separator, <220> field identifier
L:498 M:283 W: Missing Blank Line separator, <220> field identifier
L:509 M:254 E: No. of Bases conflict, this line has no nucleotides.